

Michael Gill

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

470
papers

59,150
citations

100
h-index

235
g-index

494
ext. papers

70,626
ext. citations

8.6
avg, IF

7.99
L-index

#	Paper	IF	Citations
470	Biological insights from 108 schizophrenia-associated genetic loci. <i>Nature</i> , 2014 , 511, 421-7	50.4	5249
469	Common polygenic variation contributes to risk of schizophrenia and bipolar disorder. <i>Nature</i> , 2009 , 460, 748-52	50.4	3568
468	Meta-analysis of 74,046 individuals identifies 11 new susceptibility loci for Alzheimer's disease. <i>Nature Genetics</i> , 2013 , 45, 1452-8	36.3	2714
467	Genome-wide association study identifies variants at CLU and PICALM associated with Alzheimer's disease. <i>Nature Genetics</i> , 2009 , 41, 1088-93	36.3	2018
466	Genetic relationship between five psychiatric disorders estimated from genome-wide SNPs. <i>Nature Genetics</i> , 2013 , 45, 984-94	36.3	1628
465	Synaptic, transcriptional and chromatin genes disrupted in autism. <i>Nature</i> , 2014 , 515, 209-15	50.4	1581
464	Genome-wide association study identifies five new schizophrenia loci. <i>Nature Genetics</i> , 2011 , 43, 969-76	36.3	1508
463	Functional impact of global rare copy number variation in autism spectrum disorders. <i>Nature</i> , 2010 , 466, 368-72	50.4	1499
462	Common variants at ABCA7, MS4A6A/MS4A4E, EPHA1, CD33 and CD2AP are associated with Alzheimer's disease. <i>Nature Genetics</i> , 2011 , 43, 429-35	36.3	1421
461	Genome-wide association analyses identify 44 risk variants and refine the genetic architecture of major depression. <i>Nature Genetics</i> , 2018 , 50, 668-681	36.3	1301
460	Genome-wide association analysis identifies 13 new risk loci for schizophrenia. <i>Nature Genetics</i> , 2013 , 45, 1150-9	36.3	1153
459	Large-scale genome-wide association analysis of bipolar disorder identifies a new susceptibility locus near ODZ4. <i>Nature Genetics</i> , 2011 , 43, 977-83	36.3	1094
458	Collaborative genome-wide association analysis supports a role for ANK3 and CACNA1C in bipolar disorder. <i>Nature Genetics</i> , 2008 , 40, 1056-8	36.3	949
457	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates A β tau, immunity and lipid processing. <i>Nature Genetics</i> , 2019 , 51, 414-430	36.3	917
456	Identification of loci associated with schizophrenia by genome-wide association and follow-up. <i>Nature Genetics</i> , 2008 , 40, 1053-5	36.3	877
455	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019 , 51, 793-803	36.3	662
454	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. <i>American Journal of Human Genetics</i> , 2015 , 97, 576-92	11	649

453	Convergence of genes and cellular pathways dysregulated in autism spectrum disorders. <i>American Journal of Human Genetics</i> , 2014 , 94, 677-94	11	635
452	Recurrent rearrangements of chromosome 1q21.1 and variable pediatric phenotypes. <i>New England Journal of Medicine</i> , 2008 , 359, 1685-99	59.2	587
451	Large-Scale Exome Sequencing Study Implicates Both Developmental and Functional Changes in the Neurobiology of Autism. <i>Cell</i> , 2020 , 180, 568-584.e23	56.2	578
450	Microduplications of 16p11.2 are associated with schizophrenia. <i>Nature Genetics</i> , 2009 , 41, 1223-7	36.3	550
449	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. <i>Nature Genetics</i> , 2017 , 49, 27-35	36.3	530
448	Rare coding variants in <i>PLCG2</i> , <i>ABI3</i> , and <i>TREM2</i> implicate microglial-mediated innate immunity in Alzheimer's disease. <i>Nature Genetics</i> , 2017 , 49, 1373-1384	36.3	508
447	Identification of common variants associated with human hippocampal and intracranial volumes. <i>Nature Genetics</i> , 2012 , 44, 552-61	36.3	498
446	Genome-wide association meta-analysis in 269,867 individuals identifies new genetic and functional links to intelligence. <i>Nature Genetics</i> , 2018 , 50, 912-919	36.3	475
445	A genome-wide scan for common alleles affecting risk for autism. <i>Human Molecular Genetics</i> , 2010 , 19, 4072-82	5.6	443
444	The analysis of 51 genes in DSM-IV combined type attention deficit hyperactivity disorder: association signals in <i>DRD4</i> , <i>DAT1</i> and 16 other genes. <i>Molecular Psychiatry</i> , 2006 , 11, 934-53	15.1	439
443	Confirmation of association between attention deficit hyperactivity disorder and a dopamine transporter polymorphism. <i>Molecular Psychiatry</i> , 1997 , 2, 311-3	15.1	409
442	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , 2019 , 179, 1469-1482.e11	56.2	402
441	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. <i>Cell</i> , 2018 , 173, 1705-1715.e16	56.2	360
440	Meta-analysis of genome-wide association studies of attention-deficit/hyperactivity disorder. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2010 , 49, 884-97	7.2	357
439	Genome scan meta-analysis of schizophrenia and bipolar disorder, part III: Bipolar disorder. <i>American Journal of Human Genetics</i> , 2003 , 73, 49-62	11	353
438	Mapping susceptibility loci in attention deficit hyperactivity disorder: preferential transmission of parental alleles at <i>DAT1</i> , <i>DBH</i> and <i>DRD5</i> to affected children. <i>Molecular Psychiatry</i> , 1999 , 4, 192-6	15.1	347
437	Widespread white matter microstructural differences in schizophrenia across 4322 individuals: results from the ENIGMA Schizophrenia DTI Working Group. <i>Molecular Psychiatry</i> , 2018 , 23, 1261-1269	15.1	324
436	Genome-wide association scan of quantitative traits for attention deficit hyperactivity disorder identifies novel associations and confirms candidate gene associations. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008 , 147B, 1345-54	3.5	299

435	Individual common variants exert weak effects on the risk for autism spectrum disorders. <i>Human Molecular Genetics</i> , 2012 , 21, 4781-92	5.6	279
434	Genetic evidence implicates the immune system and cholesterol metabolism in the aetiology of Alzheimer's disease. <i>PLoS ONE</i> , 2010 , 5, e13950	3.7	276
433	De novo mutations in schizophrenia implicate chromatin remodeling and support a genetic overlap with autism and intellectual disability. <i>Molecular Psychiatry</i> , 2014 , 19, 652-8	15.1	263
432	Duplications of the neuropeptide receptor gene VIPR2 confer significant risk for schizophrenia. <i>Nature</i> , 2011 , 471, 499-503	50.4	257
431	Study of 300,486 individuals identifies 148 independent genetic loci influencing general cognitive function. <i>Nature Communications</i> , 2018 , 9, 2098	17.4	254
430	High frequencies of de novo CNVs in bipolar disorder and schizophrenia. <i>Neuron</i> , 2011 , 72, 951-63	13.9	240
429	Common polygenic variation enhances risk prediction for Alzheimer's disease. <i>Brain</i> , 2015 , 138, 3673-84	11.2	227
428	Fine mapping of ZNF804A and genome-wide significant evidence for its involvement in schizophrenia and bipolar disorder. <i>Molecular Psychiatry</i> , 2011 , 16, 429-41	15.1	221
427	Evidence for novel susceptibility genes for late-onset Alzheimer's disease from a genome-wide association study of putative functional variants. <i>Human Molecular Genetics</i> , 2007 , 16, 865-73	5.6	221
426	Emotional lability in children and adolescents with attention deficit/hyperactivity disorder (ADHD): clinical correlates and familial prevalence. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2010 , 51, 915-23	7.9	214
425	Association between schizophrenia and T102C polymorphism of the 5-hydroxytryptamine type 2a-receptor gene. European Multicentre Association Study of Schizophrenia (EMASS) Group. <i>Lancet, The</i> , 1996 , 347, 1294-6	4.0	214
424	Response variability in attention deficit hyperactivity disorder: evidence for neuropsychological heterogeneity. <i>Neuropsychologia</i> , 2007 , 45, 630-8	3.2	204
423	Genome-wide association scan of attention deficit hyperactivity disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008 , 147B, 1337-44	3.5	201
422	Genome-wide analysis of copy number variants in attention deficit hyperactivity disorder: the role of rare variants and duplications at 15q13.3. <i>American Journal of Psychiatry</i> , 2012 , 169, 195-204	11.9	195
421	Dissociation in performance of children with ADHD and high-functioning autism on a task of sustained attention. <i>Neuropsychologia</i> , 2007 , 45, 2234-45	3.2	193
420	Combined analysis from eleven linkage studies of bipolar disorder provides strong evidence of susceptibility loci on chromosomes 6q and 8q. <i>American Journal of Human Genetics</i> , 2005 , 77, 582-95	11	192
419	Genome-wide significant associations in schizophrenia to ITIH3/4, CACNA1C and SDCCAG8, and extensive replication of associations reported by the Schizophrenia PGC. <i>Molecular Psychiatry</i> , 2013 , 18, 708-12	15.1	184
418	A combined analysis of D22S278 marker alleles in affected sib-pairs: support for a susceptibility locus for schizophrenia at chromosome 22q12. Schizophrenia Collaborative Linkage Group (Chromosome 22). <i>American Journal of Medical Genetics Part A</i> , 1996 , 67, 40-5		180

417	A novel Alzheimer disease locus located near the gene encoding tau protein. <i>Molecular Psychiatry</i> , 2016 , 21, 108-17	15.1	175
416	Molecular pathways involved in neuronal cell adhesion and membrane scaffolding contribute to schizophrenia and bipolar disorder susceptibility. <i>Molecular Psychiatry</i> , 2011 , 16, 286-92	15.1	175
415	Updated European Consensus Statement on diagnosis and treatment of adult ADHD. <i>European Psychiatry</i> , 2019 , 56, 14-34	6	170
414	Dissecting the attention deficit hyperactivity disorder (ADHD) phenotype: sustained attention, response variability and spatial attentional asymmetries in relation to dopamine transporter (DAT1) genotype. <i>Neuropsychologia</i> , 2005 , 43, 1847-57	3.2	169
413	Genomewide linkage scan in schizoaffective disorder: significant evidence for linkage at 1q42 close to DISC1, and suggestive evidence at 22q11 and 19p13. <i>Archives of General Psychiatry</i> , 2005 , 62, 1081-8		164
412	Joint analysis of psychiatric disorders increases accuracy of risk prediction for schizophrenia, bipolar disorder, and major depressive disorder. <i>American Journal of Human Genetics</i> , 2015 , 96, 283-94	11	161
411	Autism symptoms in Attention-Deficit/Hyperactivity Disorder: a familial trait which correlates with conduct, oppositional defiant, language and motor disorders. <i>Journal of Autism and Developmental Disorders</i> , 2009 , 39, 197-209	4.6	161
410	Identification in 2 independent samples of a novel schizophrenia risk haplotype of the dystrobrevin binding protein gene (DTNBP1). <i>Archives of General Psychiatry</i> , 2004 , 61, 336-44		155
409	A novel approach of homozygous haplotype sharing identifies candidate genes in autism spectrum disorder. <i>Human Genetics</i> , 2012 , 131, 565-79	6.3	150
408	Delay and reward choice in ADHD: an experimental test of the role of delay aversion. <i>Neuropsychology</i> , 2009 , 23, 367-80	3.8	149
407	Joint analysis of the DRD5 marker concludes association with attention-deficit/hyperactivity disorder confined to the predominantly inattentive and combined subtypes. <i>American Journal of Human Genetics</i> , 2004 , 74, 348-56	11	148
406	Investigating the contribution of common genetic variants to the risk and pathogenesis of ADHD. <i>American Journal of Psychiatry</i> , 2012 , 169, 186-94	11.9	147
405	Convergent genetic and expression data implicate immunity in Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2015 , 11, 658-71	1.2	146
404	Evidence that interaction between neuregulin 1 and its receptor erbB4 increases susceptibility to schizophrenia. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2006 , 141B, 96-103	3.5	143
403	Molecular genetic evidence for overlap between general cognitive ability and risk for schizophrenia: a report from the Cognitive Genomics consortium (COGENT). <i>Molecular Psychiatry</i> , 2014 , 19, 168-74	15.1	142
402	Detecting subtle facial emotion recognition deficits in high-functioning Autism using dynamic stimuli of varying intensities. <i>Neuropsychologia</i> , 2010 , 48, 2777-81	3.2	134
401	Whole genome linkage scan of recurrent depressive disorder from the depression network study. <i>Human Molecular Genetics</i> , 2005 , 14, 3337-45	5.6	133
400	Serotonergic system and attention deficit hyperactivity disorder (ADHD): a potential susceptibility locus at the 5-HT(1B) receptor gene in 273 nuclear families from a multi-centre sample. <i>Molecular Psychiatry</i> , 2002 , 7, 718-25	15.1	132

399	Excess of rare novel loss-of-function variants in synaptic genes in schizophrenia and autism spectrum disorders. <i>Molecular Psychiatry</i> , 2014 , 19, 872-9	15.1	131
398	Case-control genome-wide association study of attention-deficit/hyperactivity disorder. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2010 , 49, 906-20	7.2	131
397	Meta-analysis of genome-wide linkage scans of attention deficit hyperactivity disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008 , 147B, 1392-8	3.5	131
396	Evidence that duplications of 22q11.2 protect against schizophrenia. <i>Molecular Psychiatry</i> , 2014 , 19, 37-40	15.1	130
395	Separation of cognitive impairments in attention-deficit/hyperactivity disorder into 2 familial factors. <i>Archives of General Psychiatry</i> , 2010 , 67, 1159-67		130
394	Evidence that variation at the serotonin transporter gene influences susceptibility to attention deficit hyperactivity disorder (ADHD): analysis and pooled analysis. <i>Molecular Psychiatry</i> , 2002 , 7, 908-12	15.1	130
393	Synaptosomal-associated protein 25 (SNAP-25) and attention deficit hyperactivity disorder (ADHD): evidence of linkage and association in the Irish population. <i>Molecular Psychiatry</i> , 2002 , 7, 913-7	15.1	128
392	The influence of serotonin- and other genes on impulsive behavioral aggression and cognitive impulsivity in children with attention-deficit/hyperactivity disorder (ADHD): Findings from a family-based association test (FBAT) analysis. <i>Behavioral and Brain Functions</i> , 2008 , 4, 48	4.1	127
391	Dopaminergic system genes in ADHD: toward a biological hypothesis. <i>Neuropsychopharmacology</i> , 2002 , 27, 607-19	8.7	127
390	Dementia in people with Down's syndrome. <i>International Journal of Geriatric Psychiatry</i> , 2001 , 16, 1168-74	4.9	127
389	Confirmation and refinement of an 'at-risk' haplotype for schizophrenia suggests the EST cluster, Hs.97362, as a potential susceptibility gene at the Neuregulin-1 locus. <i>Molecular Psychiatry</i> , 2004 , 9, 208-13	15.1	126
388	Allelic association between a Ser-9-Gly polymorphism in the dopamine D3 receptor gene and schizophrenia. <i>Human Genetics</i> , 1996 , 97, 714-9	6.3	126
387	Follow-up of a report of a potential linkage for schizophrenia on chromosome 22q12-q13.1: Part 2. <i>American Journal of Medical Genetics Part A</i> , 1994 , 54, 44-50		124
386	GWAS meta-analysis reveals novel loci and genetic correlates for general cognitive function: a report from the COGENT consortium. <i>Molecular Psychiatry</i> , 2017 , 22, 336-345	15.1	123
385	Psychosis susceptibility gene ZNF804A and cognitive performance in schizophrenia. <i>Archives of General Psychiatry</i> , 2010 , 67, 692-700		120
384	Association of the 480 bp DAT1 allele with methylphenidate response in a sample of Irish children with ADHD. <i>American Journal of Medical Genetics Part A</i> , 2003 , 121B, 50-4		120
383	Linkage disequilibrium mapping at DAT1, DRD5 and DBH narrows the search for ADHD susceptibility alleles at these loci. <i>Molecular Psychiatry</i> , 2003 , 8, 299-308	15.1	119
382	The SNP ratio test: pathway analysis of genome-wide association datasets. <i>Bioinformatics</i> , 2009 , 25, 2767-3	7.3	116

381	Confirmation that a specific haplotype of the dopamine transporter gene is associated with combined-type ADHD. <i>American Journal of Psychiatry</i> , 2007 , 164, 674-7	11.9	115
380	Confirming RGS4 as a susceptibility gene for schizophrenia. <i>American Journal of Medical Genetics Part A</i> , 2004 , 125B, 50-3		114
379	Dopamine D4 receptor subtypes and response to clozapine. <i>Lancet, The</i> , 1993 , 341, 116	40	114
378	DSM-IV combined type ADHD shows familial association with sibling trait scores: a sampling strategy for QTL linkage. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008 , 147B, 1450-60	3.5	113
377	Four components describe behavioral symptoms in 1,120 individuals with late-onset Alzheimer's disease. <i>Journal of the American Geriatrics Society</i> , 2006 , 54, 1348-54	5.6	111
376	DNA methylation of the serotonin transporter gene in peripheral cells and stress-related changes in hippocampal volume: a study in depressed patients and healthy controls. <i>PLoS ONE</i> , 2015 , 10, e0119067	2.7	110
375	Impaired conflict resolution and alerting in children with ADHD: evidence from the Attention Network Task (ANT). <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2008 , 49, 1339-47	7.9	106
374	Tryptophan hydroxylase 2 (TPH2) gene variants associated with ADHD. <i>Molecular Psychiatry</i> , 2005 , 10, 944-9	15.1	104
373	Association of the paternally transmitted copy of common Valine allele of the Val66Met polymorphism of the brain-derived neurotrophic factor (BDNF) gene with susceptibility to ADHD. <i>Molecular Psychiatry</i> , 2005 , 10, 939-43	15.1	102
372	Variance in neurocognitive performance is associated with dysbindin-1 in schizophrenia: a preliminary study. <i>Neuropsychologia</i> , 2007 , 45, 454-8	3.2	101
371	The gene for Darier's disease maps to chromosome 12q23-q24.1. <i>Human Molecular Genetics</i> , 1993 , 2, 1941-3	5.6	100
370	Linkage studies of bipolar disorder in the region of the Darier's disease gene on chromosome 12q23-24.1. <i>American Journal of Medical Genetics Part A</i> , 1995 , 60, 94-102		98
369	No evidence for an association of affective disorders with high- or low-activity allele of catechol-o-methyltransferase gene. <i>Biological Psychiatry</i> , 1997 , 42, 282-5	7.9	97
368	Development of strategies for SNP detection in RNA-seq data: application to lymphoblastoid cell lines and evaluation using 1000 Genomes data. <i>PLoS ONE</i> , 2013 , 8, e58815	3.7	94
367	Conduct disorder and ADHD: evaluation of conduct problems as a categorical and quantitative trait in the international multicentre ADHD genetics study. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008 , 147B, 1369-78	3.5	93
366	A Genetic Investigation of Sex Bias in the Prevalence of Attention-Deficit/Hyperactivity Disorder. <i>Biological Psychiatry</i> , 2018 , 83, 1044-1053	7.9	93
365	Genome-wide association scan of the time to onset of attention deficit hyperactivity disorder. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008 , 147B, 1355-8	3.5	92
364	Preferential transmission of paternal alleles at risk genes in attention-deficit/hyperactivity disorder. <i>American Journal of Human Genetics</i> , 2005 , 77, 958-65	11	92

363	Association of DRD4 in children with ADHD and comorbid conduct problems. <i>American Journal of Medical Genetics Part A</i> , 2002 , 114, 150-3		92
362	Increased BDNF levels and NTRK2 gene association suggest a disruption of BDNF/TrkB signaling in autism. <i>Genes, Brain and Behavior</i> , 2010 , 9, 841-8	3.6	91
361	Genome-wide association study of Alzheimer's disease with psychotic symptoms. <i>Molecular Psychiatry</i> , 2012 , 17, 1316-27	15.1	90
360	Gene-wide analysis detects two new susceptibility genes for Alzheimer's disease. <i>PLoS ONE</i> , 2014 , 9, e94661	3.7	90
359	Human subcortical brain asymmetries in 15,847 people worldwide reveal effects of age and sex. <i>Brain Imaging and Behavior</i> , 2017 , 11, 1497-1514	4.1	87
358	A review of gene linkage, association and expression studies in autism and an assessment of convergent evidence. <i>International Journal of Developmental Neuroscience</i> , 2007 , 25, 69-85	2.7	85
357	Genomewide association scan of suicidal thoughts and behaviour in major depression. <i>PLoS ONE</i> , 2011 , 6, e20690	3.7	83
356	Health co-morbidities in ageing persons with Down syndrome and Alzheimer's dementia. <i>Journal of Intellectual Disability Research</i> , 2005 , 49, 560-6	3.2	82
355	Attention network hypoconnectivity with default and affective network hyperconnectivity in adults diagnosed with attention-deficit/hyperactivity disorder in childhood. <i>JAMA Psychiatry</i> , 2013 , 70, 1329-37	14.5	81
354	Evaluation of a susceptibility gene for schizophrenia: genotype based meta-analysis of RGS4 polymorphisms from thirteen independent samples. <i>Biological Psychiatry</i> , 2006 , 60, 152-62	7.9	80
353	Familiality of symptom dimensions in depression. <i>Archives of General Psychiatry</i> , 2004 , 61, 468-74		80
352	No association of the dopamine DRD4 receptor (DRD4) gene polymorphism with attention deficit hyperactivity disorder (ADHD) in the Irish population. <i>American Journal of Medical Genetics Part A</i> , 2000 , 96, 268-72		80
351	Imprinting and anticipation. Are they relevant to genetic studies of schizophrenia?. <i>British Journal of Psychiatry</i> , 1994 , 164, 619-24	5.4	80
350	Oxytocin receptor (OXTR) does not play a major role in the aetiology of autism: genetic and molecular studies. <i>Neuroscience Letters</i> , 2010 , 474, 163-167	3.3	79
349	Association between dopamine transporter (DAT1) genotype, left-sided inattention, and an enhanced response to methylphenidate in attention-deficit hyperactivity disorder. <i>Neuropsychopharmacology</i> , 2005 , 30, 2290-7	8.7	79
348	Shared polygenic contribution between childhood attention-deficit hyperactivity disorder and adult schizophrenia. <i>British Journal of Psychiatry</i> , 2013 , 203, 107-11	5.4	78
347	Serotonin transporter gene and autism: a haplotype analysis in an Irish autistic population. <i>Molecular Psychiatry</i> , 2004 , 9, 587-93	15.1	78
346	Evidence for a genetic association between alleles of monoamine oxidase A gene and bipolar affective disorder. <i>American Journal of Medical Genetics Part A</i> , 1995 , 60, 325-31		78

345	Attention deficit/hyperactivity disorder-derived coding variation in the dopamine transporter disrupts microdomain targeting and trafficking regulation. <i>Journal of Neuroscience</i> , 2012 , 32, 5385-97	6.6	77
344	Cigarette smoking and psychotic symptoms in bipolar affective disorder. <i>British Journal of Psychiatry</i> , 2001 , 179, 35-8	5.4	77
343	Linkage studies on chromosome 22 in familial schizophrenia. <i>American Journal of Medical Genetics Part A</i> , 1995 , 60, 139-46		77
342	DRD4 gene variants and sustained attention in attention deficit hyperactivity disorder (ADHD): effects of associated alleles at the VNTR and -521 SNP. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2005 , 136B, 81-6	3.5	75
341	The methionine allele of the COMT polymorphism impairs prefrontal cognition in children and adolescents with ADHD. <i>Experimental Brain Research</i> , 2005 , 163, 352-60	2.3	75
340	Biological overlap of attention-deficit/hyperactivity disorder and autism spectrum disorder: evidence from copy number variants. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2014 , 53, 761-70.e26	7.2	74
339	ZNF804A risk allele is associated with relatively intact gray matter volume in patients with schizophrenia. <i>NeuroImage</i> , 2011 , 54, 2132-7	7.9	74
338	The cognitive genetics of attention deficit hyperactivity disorder (ADHD): sustained attention as a candidate phenotype. <i>Cortex</i> , 2006 , 42, 838-45	3.8	74
337	No evidence for association of the dysbindin gene [DTNBP1] with schizophrenia in an Irish population-based study. <i>Schizophrenia Research</i> , 2003 , 60, 167-72	3.6	74
336	Confirmation of association between autism and the mitochondrial aspartate/glutamate carrier SLC25A12 gene on chromosome 2q31. <i>American Journal of Psychiatry</i> , 2005 , 162, 2182-4	11.9	74
335	Bipolar affective puerperal psychosis: genome-wide significant evidence for linkage to chromosome 16. <i>American Journal of Psychiatry</i> , 2007 , 164, 1099-104	11.9	73
334	Schizophrenia: genetics and the maternal immune response to viral infection. <i>American Journal of Medical Genetics Part A</i> , 1993 , 48, 40-6		73
333	Single-Nucleotide Polymorphism of the FKBP5 Gene and Childhood Maltreatment as Predictors of Structural Changes in Brain Areas Involved in Emotional Processing in Depression. <i>Neuropsychopharmacology</i> , 2016 , 41, 487-97	8.7	72
332	Large-Scale Cognitive GWAS Meta-Analysis Reveals Tissue-Specific Neural Expression and Potential Nootropic Drug Targets. <i>Cell Reports</i> , 2017 , 21, 2597-2613	10.6	71
331	Neuropsychological correlates of emotional lability in children with ADHD. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2012 , 53, 1139-48	7.9	71
330	Brain-derived neurotrophic factor Val66Met polymorphism and early life adversity affect hippocampal volume. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2013 , 162B, 183-90	3.5	71
329	Does parental expressed emotion moderate genetic effects in ADHD? An exploration using a genome wide association scan. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008 , 147B, 1359-68	3.5	71
328	Genetic relationships between suicide attempts, suicidal ideation and major psychiatric disorders: a genome-wide association and polygenic scoring study. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2014 , 165B, 428-37	3.5	70

327	Noradrenergic genotype predicts lapses in sustained attention. <i>Neuropsychologia</i> , 2009 , 47, 591-4	3.2	69
326	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. <i>Biological Psychiatry</i> , 2017 , 82, 322-329	7.9	68
325	Common variant at 16p11.2 conferring risk of psychosis. <i>Molecular Psychiatry</i> , 2014 , 19, 108-14	15.1	67
324	Agony and ecstasy: a review of MDMA effects and toxicity. <i>European Psychiatry</i> , 2000 , 15, 287-94	6	67
323	Familiality of symptom dimensions in schizophrenia. <i>Schizophrenia Research</i> , 2001 , 47, 223-32	3.6	67
322	DRD2 Ser311/Cys311 polymorphism in schizophrenia. <i>Lancet, The</i> , 1994 , 343, 1044-1046	4.0	67
321	A review of neuropsychological and neuroimaging research in autistic spectrum disorders: Attention, inhibition and cognitive flexibility. <i>Research in Autism Spectrum Disorders</i> , 2008 , 2, 1-16	3	66
320	Linkage analysis of manic depression in an Irish family using H-ras 1 and INS DNA markers. <i>Journal of Medical Genetics</i> , 1988 , 25, 634-5	5.8	65
319	European Multicentre Association Study of Schizophrenia: a study of the DRD2 Ser311Cys and DRD3 Ser9Gly polymorphisms. <i>American Journal of Medical Genetics Part A</i> , 1998 , 81, 24-8		64
318	Evidence that genetic variation in the oxytocin receptor (OXTR) gene influences social cognition in ADHD. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2010 , 34, 697-702	5.5	63
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316	The Wellcome trust UK-Irish bipolar affective disorder sibling-pair genome screen: first stage report. <i>Molecular Psychiatry</i> , 2002 , 7, 189-200	15.1	63
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